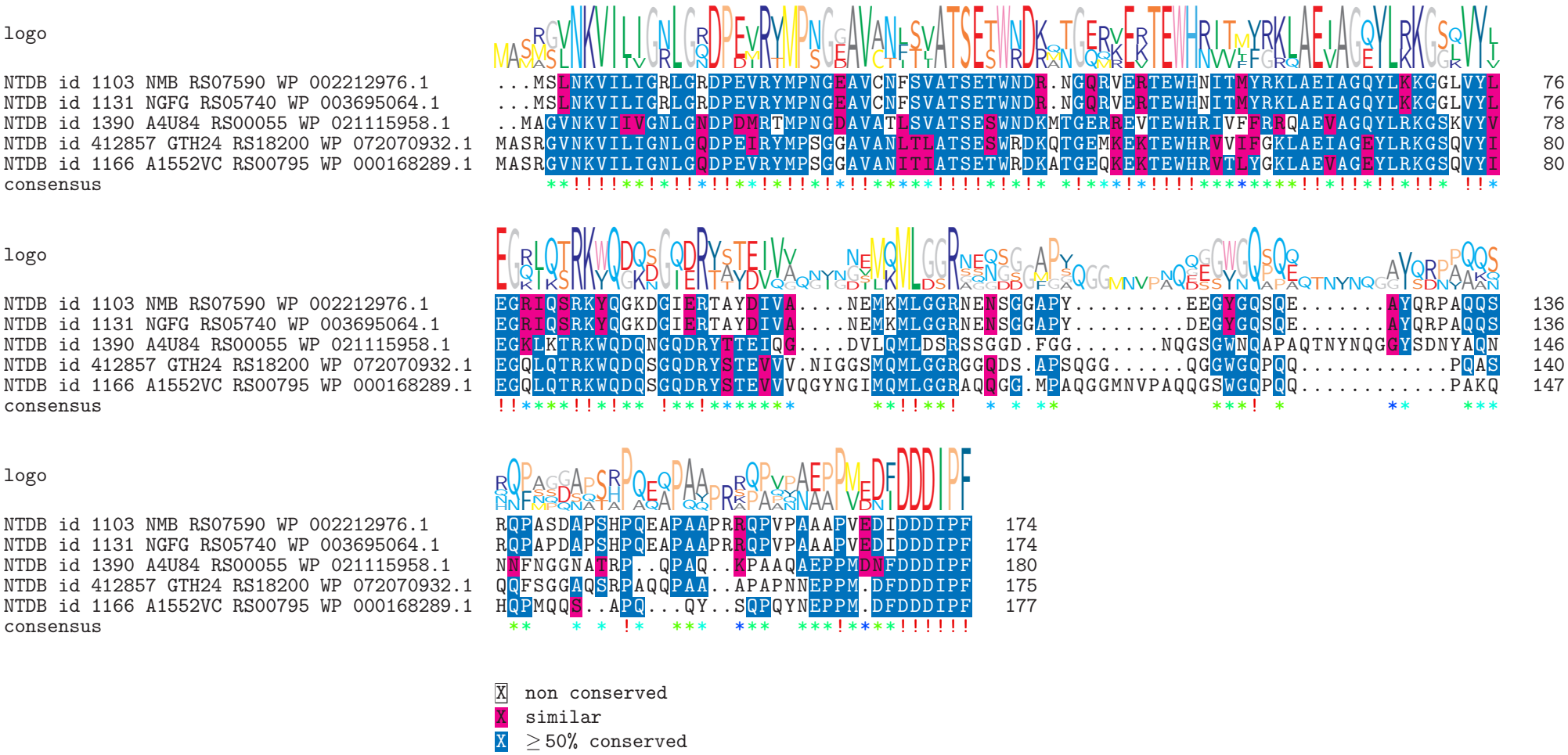
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
NTDB id 412857 GTH24 RS18200 WP 072070932.1 MASRGVNKVILIGNLGQDPEIRYMPSGGAVANLTLATSESWRDKQTGEMKEKTEWHRVVIFGKLAEIAGEYLRKGSQVYI 80
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPY.........EEGYGQSQE.......AYQRPAQQS 136
NTDB id 1131 NGFG RS05740 WP 003695064.1 EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPY.........DEGYGQSQE.......AYQRPAQQS 136
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGG.......NQGSGWNQAPAQTNYNQGGYSDNYAQN 146
NTDB id 412857 GTH24 RS18200 WP 072070932.1 EGQLQTRKWQDQSGQDRYSTEVVV.NIGGSMQMLGGRGGQDS.APSQGG......QGGWGQPQQ............PQAS 140
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGMNVPAQQGSWGQPQQ............PAKQ 147
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 RQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1131 NGFG RS05740 WP 003695064.1 RQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1390 A4U84 RS00055 WP 021115958.1 NNFNGGNATRP..QPAQ..KPAAQAEPPMDNFDDDIPF 180
NTDB id 412857 GTH24 RS18200 WP 072070932.1 QQFSGGAQSRPAQQPAA..APAPNNEPPM.DFDDDIPF 175
NTDB id 1166 A1552VC RS00795 WP 000168289.1 HQPMQQS..APQ...QY..SQPQYNEPPM.DFDDDIPF 177
consensus
X non conserved
X similar
X ≥ 50% conserved